

RESULT 4
 AP002569/c
 LOCUS AP002569 168764 bp DNA linear BCT 07-MAR-2001
 DEFINITION *Escherichia coli* O157:H7 DNA, complete genome, section 20/20.
 ACCESSION AP002569 BA000007
 VERSION AP002569.1 GI:13364704
 KEYWORDS .
 SOURCE *Escherichia coli* O157:H7
 ORGANISM *Escherichia coli* O157:H7
 Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
 Enterobacteriaceae; *Escherichia*.
 REFERENCE 1 (sites)
 AUTHORS Makino,K., Yokoyama,K., Kubota,Y., Yutsudo,C.H., Kimura,S.,
 Kurokawa,K., Ishii,K., Hattori,M., Tatsuno,I., Abe,H., Iida,T.,
 Yamamoto,K., Ohnishi,M., Hayashi,T., Yasunaga,T., Honda,T.,
 Sasakawa,C. and Shinagawa,H.
 TITLE Complete nucleotide sequence of the prophage VT2-Sakai carrying the
 verotoxin 2 genes of the enterohemorrhagic *Escherichia coli* O157:H7
 derived from the Sakai outbreak
 JOURNAL Genes Genet. Syst. 74 (5), 227-239 (1999)
 MEDLINE 20198780
 PUBMED 10734605
 REFERENCE 2 (sites)
 AUTHORS Ohnishi,M., Murata,T., Nakayama,K., Kuhara,S., Hattori,M.,
 Kurokawa,K., Yasunaga,T., Yokoyama,K., Makino,K., Shinagawa,H. and
 Hayashi,T.
 TITLE Comparative analysis of the whole set of rRNA operons between an
 enterohemorrhagic *Escherichia coli* O157:H7 Sakai strain and an
Escherichia coli K-12 strain MG1655
 JOURNAL Syst. Appl. Microbiol. 23 (3), 315-324 (2000)
 MEDLINE 20557356
 PUBMED 11108008
 REFERENCE 3 (sites)
 AUTHORS Yokoyama,K., Makino,K., Kubota,Y., Watanabe,M., Kimura,S.,
 Yutsudo,C.H., Kurokawa,K., Ishii,K., Hattori,M., Abe,H., Iida,T.,
 Yamamoto,K., Hayashi,T., Yasunaga,T., Honda,T., Sasakawa,C. and
 Shinagawa,H.
 TITLE Complete nucleotide sequence of the prophage VT1-Sakai carrying the
 Shiga toxin 1 genes of the enterohemorrhagic *Escherichia coli*
 O157:H7 strain derived from the Sakai outbreak
 JOURNAL Gene 258 (1-2), 127-139 (2000)
 MEDLINE 20564182
 PUBMED 11111050
 REFERENCE 4 (sites)
 AUTHORS Hayashi,T., Makino,K., Ohnishi,M., Kurokawa,K., Ishii,K.,
 Yokoyama,K., Han,C.-G., Ohtsubo,E., Nakayama,K., Murata,T.,
 Tanaka,M., Tobe,T., Iida,T., Takami,H., Honda,T., Sasakawa,C.,
 Ogasawara,N., Yasunaga,T., Kuhara,S., Shiba,T., Hattori,M. and
 Shinagawa,H.
 TITLE Complete genome sequence of enterohemorrhagic *Escherichia coli*
 O157:H7 and genomic comparison with a laboratory strain K-12
 JOURNAL DNA Res. 8 (1), 11-22 (2001)
 MEDLINE 21156231
 PUBMED 11258796
 REFERENCE 5 (bases 1 to 168764)
 AUTHORS Ohnishi,M., Kurokawa,K., Makino,K., Yasunaga,T., Shinagawa,H. and
 Hayashi,T.